## SEQUENCE LISTING

<110> Raju, Jeyaseelan

<120> NOVEL CARK PROTEIN AND NUCLEIC ACID MOLECULES AND USES THEREFOR

<130> MNI-068CP2

<140> US 09/947,199

<141> 2001-09-05

<150> US 60/111,938

<151> 1998-12-11

<150> US 09/291,839

<151> 1999-04-14

<150> US 09/458,457

<151> 1999-12-10

<160> 9

<170> PatentIn Ver. 2.0

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<211> 3025

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<213> Homo sapiens

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<221> CDS

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gtc agt gaa tca tat gtt atc aca ata gaa aga tta gaa gat gac ctg 152 Val Ser Glu Ser Tyr Val Ile Thr Ile Glu Arg Leu Glu Asp Asp Leu 20 25 30 35

cag atc aag gaa aaa gaa ctg aca gaa cta agg aat ata ttt ggc tct 200 Gln Ile Lys Glu Leu Thr Glu Leu Arg Asn Ile Phe Gly Ser

gat gaa gcc ttc agt aaa gtc aat tta aat tac cgc act gaa aat ggg 248 Asp Glu Ala Phe Ser Lys Val Asn Leu Asn Tyr Arg Thr Glu Asn Gly 55 60 65

ctg tct cta ctt cat tta tgt tgc att tgt gga ggc aag aaa tca cat 296

Leu	Ser	Leu 70	Leu	His	Leu	Cys	Cys 75	Ile	Cys	Gly	Gly	Lys 80	Lys	Ser	His	
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										tac Tyr 110						392
										gat Asp						440
										aca Thr		_				488
	_	_	_		_					gct Ala		_				536
										gca Ala						584
										ggt Gly 190					_	632
										cta Leu						680
										gaa Glu						728
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										gga Gly 270						872
		_						_	_	gcc Ala	_	_				920
ata	tca	gga	aca	gaa	agt	ctg	act	aag	gaa	aac	atc	ttc	agt	gaa	aca	968

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	tcc Ser					_	_				_	_	_			1736
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_	aca Thr	_					_	_	_		_					1832
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	aac Asn 725															2264
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Leu Ser Pro Ser Ser Ser Ser Asp Cys Leu Val Asn Arg Gly Gly Pro 740 745 750 755
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gct cta aat gca agg tcc tat gct gct ttg tcc caa agt gct gga caa 2408 Ala Leu Asn Ala Arg Ser Tyr Ala Ala Leu Ser Gln Ser Ala Gly Gln 775 780 785
tat tcc tct caa ggt ctg tct ttg gag gag atg aaa aga agt ctt caa 2456 Tyr Ser Ser Gln Gly Leu Ser Leu Glu Glu Met Lys Arg Ser Leu Gln 790 795 800
tac aca ccc att gac aaa tat ggc tat gta tcc gat ccc atg agc tca 2504 Tyr Thr Pro Ile Asp Lys Tyr Gly Tyr Val Ser Asp Pro Met Ser Ser 805 810 815
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Asp Asp Leu Gln Ile Lys Glu Lys Glu Leu Thr Glu Leu Arg Asn Ile 35 40 45

Phe Gly Ser Asp Glu Ala Phe Ser Lys Val Asn Leu Asn Tyr Arg Thr

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Lys	Ser	His	Ile	Arg 85	Thr	Leu	Met	Leu	Lys 90	Gly	Leu	Arg	Pro	Ser 95	Arg
Leu	Thr	Arg	Asn 100	Gly	Phe	Thr	Ala	Leu 105	His	Leu	Ala	Val	Tyr 110	Lys	Asp
Asn	Ala	Glu 115	Leu	Ile	Thr	Ser	Leu 120	Leu	His	Ser	Gly	Ala 125	Asp	Ile	Gln
Gln	Val 130	Gly	Tyr	Gly	Gly	Leu 135	Thr	Ala	Leu	His	Ile 140	Ala	Thr	Ile	Ala
Gly 145	His	Leu	Glu	Ala	Ala 150	Asp	Val	Leu	Leu	Gln 155	His	Gly	Ala	Asn	Val 160
Asn	Ile	Gln	Asp	Ala 165	Val	Phe	Phe	Thr	Pro 170	Leu	His	Ile	Ala	Ala 175	Tyr
Tyr	Gly	His	Glu 180	Gln	Val	Thr	Arg	Leu 185	Leu	Leu	Lys	Phe	Gly 190	Ala	Asp
Val	Asn	Val 195	Ser	Gly	Glu	Val	Gly 200	Asp	Arg	Pro	Leu	His 205	Leu	Ala	Ser
Ala	Lys 210	Gly	Phe	Leu	Asn	Ile 215	Ala	Lys	Leu	Leu	Met 220	Glu	Glu	Gly	Ser
Lys 225	Ala	Asp	Val	Asn	Ala 230	Gln	Asp	Asn	Glu	Asp 235	His	Val	Pro	Leu	His 240
Phe	Сув	Ser	Arg	Phe 245	Gly	His	His	Asp	Ile 250	Val	Lys	Tyr	Leu	Leu 255	Gln
Ser	Asp	Leu	Glu 260	Val	Gln	Pro	His	Val 265	Val	Asn	Ile	Tyr	Gly 270	Asp	Thr
Pro	Leu	His 275	Leu	Ala	Суѕ	Tyr	Asn 280	Gly	Lys	Phe	Glu	Val 285	Ala	Lys	Glu
Ile	Ile 290	Gln	Ile	Ser	Gly	Thr 295	Glu	Ser	Leu	Thr	Lys 300	Glu	Asn	Ile	Phe
Ser 305	Glu	Thr	Ala	Phe	His 310	Ser	Ala	Сув	Thr	Tyr 315	Gly	Lys	Ser	Ile	Asp 320
Leu	Val	Lys	Phe	Leu 325	Leu	Asp	Gln	Asn	Val 330	Ile	Asn	Ile	Asn	His 335	Gln

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- Val Ala Cys Asp Pro Ser Arg Ser Ser Gly Glu Lys Asp Glu Gln Thr 370 375 380
- Cys Leu Met Trp Ala Tyr Glu Lys Gly His Asp Ala Ile Val Thr Leu 385 390 395 400
- Leu Lys His Tyr Lys Arg Pro Gln Asp Glu Leu Pro Cys Asn Glu Tyr 405 410 415
- Ser Gln Pro Gly Gly Asp Gly Ser Tyr Val Ser Val Pro Ser Pro Leu 420 425 430
- Gly Lys Ile Lys Ser Met Thr Lys Glu Lys Ala Asp Ile Leu Leu Leu 435 440
- Arg Ala Gly Leu Pro Ser His Phe His Leu Gln Leu Ser Glu Ile Glu 450 455 460
- Phe His Glu Ile Ile Gly Ser Gly Ser Phe Gly Lys Val Tyr Lys Gly 465 470 475 480
- Arg Cys Arg Asn Lys Ile Val Ala Ile Lys Arg Tyr Arg Ala Asn Thr 485 490 495
- Tyr Cys Ser Lys Ser Asp Val Asp Met Phe Cys Arg Glu Val Ser Ile 500 505 510
- Leu Cys Gln Leu Asn His Pro Cys Val Ile Gln Phe Val Gly Ala Cys 515 520 525
- Leu Asn Asp Pro Ser Gln Phe Ala Ile Val Thr Gln Tyr Ile Ser Gly 530 540
- Gly Ser Leu Phe Ser Leu Leu His Glu Gln Lys Arg Ile Leu Asp Leu 545 550 555 560
- Gln Ser Lys Leu Ile Ile Ala Val Asp Val Ala Lys Gly Met Glu Tyr 565 570 575
- Leu His Asn Leu Thr Gln Pro Ile Ile His Arg Asp Leu Asn Ser His 580 585 590
- Asn Ile Leu Leu Tyr Glu Asp Gly His Ala Val Val Ala Asp Phe Gly 595 600 605
- Glu Ser Arg Phe Leu Gln Ser Leu Asp Glu Asp Asn Met Thr Lys Gln 610 620
- Pro Gly Asn Leu Arg Trp Met Ala Pro Glu Val Phe Thr Gln Cys Thr 625 630 635 640
- Arg Tyr Thr Ile Lys Ala Asp Val Phe Ser Tyr Ala Leu Cys Leu Trp 645 650 655

Glu Ile Leu Thr Gly Glu Ile Pro Phe Ala His Leu Lys Pro Ala Ala 665 Ala Ala Ala Asp Met Ala Tyr His His Ile Arg Pro Pro Ile Gly Tyr Ser Ile Pro Lys Pro Ile Ser Ser Leu Leu Ile Arg Gly Trp Asn Ala 695 Cys Pro Glu Gly Arg Pro Glu Phe Ser Glu Val Val Met Lys Leu Glu 715 Glu Cys Leu Cys Asn Ile Glu Leu Met Ser Pro Ala Ser Ser Asn Ser 730 Ser Gly Ser Leu Ser Pro Ser Ser Ser Ser Asp Cys Leu Val Asn Arg 745 Gly Gly Pro Gly Arg Ser His Val Ala Ala Leu Arg Ser Arg Phe Glu Leu Glu Tyr Ala Leu Asn Ala Arg Ser Tyr Ala Ala Leu Ser Gln Ser Ala Gly Gln Tyr Ser Ser Gln Gly Leu Ser Leu Glu Glu Met Lys Arg 790 Ser Leu Gln Tyr Thr Pro Ile Asp Lys Tyr Gly Tyr Val Ser Asp Pro 810 Met Ser Ser Met His Phe His Ser Cys Arg Asn Ser Ser Ser Phe Glu

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gat gac ctg cag atc aag gaa aaa gaa ctg aca gaa cta agg aat ata 144

Asp	Asp	Leu 35	Gln	Ile	Lys	Glu	Lys 40	Glu	Leu	Thr	Glu	Leu 45	Arg	Asn	Ile	
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	aat Asn															240
	tca Ser			_				_				_			_	288
	aca Thr															336
	gca Ala	_	_				_			_		_	_		_	384
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	cac His			_	_	_		_	_				_		_	480
	att Ile															528
	gga Gly		_		-											576
~	aat Asn	_	_		_	_		_	_					_		624
_	aaa Lys 210			_			_			_	_	_	-		_	672
	gca Ala															720
	tgt Cys		_					_		-	_		_	_		768
agt	gat	ttg	gaa	gtt	caa	cct	cat	gtt	gtt	aat	atc	tat	gga	gat	acc	816

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					tgc Cys											864
					gga Gly											912
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					ctt Leu											1008
gga Gly	agg Arg	gat Asp	999 Gly 340	cac His	act Thr	gga Gly	tta Leu	cac His 345	tct Ser	gct Ala	tgc Cys	tac Tyr	cac His 350	ggt Gly	cac His	1056
			gtt		ttc Phe											1104
gtg Val	gct Ala 370	tgt Cys	gat Asp	ccc Pro	agc Ser	agg Arg 375	tct Ser	agt Ser	ggt Gly	gaa Glu	aaa Lys 380	gat Asp	gag Glu	cag Gln	aca Thr	1152
					tat Tyr 390											1200
ctg Leu	aag Lys	cat His	tat Tyr	aag Lys 405	aga Arg	cca Pro	caa Gln	gat Asp	gaa Glu 410	ttg Leu	ccc Pro	tgt Cys	aat Asn	gaa Glu 415	tat Tyr	1248
					gat Asp											1296
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aga Arg	gct Ala 450	gga Gly	ttg Leu	cct Pro	tca Ser	cat His 455	ttc Phe	cat His	ctt Leu	cag Gln	ctc Leu 460	tca Ser	gaa Glu	att Ile	gag Glu	1392
ttc Phe 465	cat His	gag Glu	att Ile	att Ile	ggc Gly 470	tca Ser	ggt Gly	tct Ser	ttt Phe	999 Gly 475	aaa Lys	gta Val	tat Tyr	aaa Lys	gga Gly 480	1440
cga	tgc	aga	aat	aaa	ata	gtg	gct	ata	aaa	cgt	tat	cga	gcc	aat	acc	1488

Arg	Cys	Arg	Asn	Lys 485	Ile	Val	Ala	Ile	Lys 490	Arg	Tyr	Arg	Ala	Asn 495	Thr	
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ttg Leu	aat Asn 530	gat Asp	ccc Pro	agc Ser	cag Gln	ttt Phe 535	gcc Ala	att Ile	gtc Val	act Thr	caa Gln 540	tac Tyr	ata Ile	tca Ser	gly aaa	1632
ggt Gly 545	tct Ser	ctg Leu	ttc Phe	tcc Ser	ctc Leu 550	ctt Leu	cat His	gag Glu	cag Gln	aag Lys 555	agg Arg	att Ile	ctt Leu	gat Asp	ttg Leu 560	1680
cag Gln	tct Ser	aaa Lys	tta Leu	att Ile 565	att Ile	gca Ala	gta Val	gat Asp	gtt Val 570	gcc Ala	aaa Lys	ggc Gly	atg Met	gag Glu 575	tac Tyr	1728
ctt Leu	cac His	aac Asn	ctg Leu 580	aca Thr	cag Gln	cca Pro	att Ile	ata Ile 585	cat His	cgt Arg	gac Asp	ttg Leu	aac Asn 590	agt Ser	cac His	1776
aat Asn	att Ile	ctt Leu 595	ctc Leu	tat Tyr	gag Glu	gat Asp	600 GJA aaa	cat His	gct Ala	gtg Val	gtg Val	gca Ala 605	gat Asp	ttt Phe	gga Gly	1824
gaa Glu	tca Ser 610	aga Arg	ttt Phe	cta Leu	cag Gln	tct Ser 615	ctg Leu	gat Asp	gaa Glu	gac Asp	aac Asn 620	atg Met	aca Thr	aaa Lys	caa Gln	1872
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gcg Ala	gca Ala	gca Ala 675	gac Asp	atg Met	gct Ala	tac Tyr	cac His 680	cac His	atc Ile	aga Arg	cct Pro	ccc Pro 685	att Ile	ggc Gly	tat Tyr	2064
tcc Ser	att Ile 690	ccc Pro	aag Lys	ccc Pro	ata Ile	tca Ser 695	tct Ser	ctg Leu	ctg Leu	ata Ile	cga Arg 700	gly aaa	tgg Trp	aac Asn	gca Ala	2112
tgt	cct	gaa	gga	aga	ccc	gaa	ttt	tct	gaa	gtt	gtc	atg	aag	tta	gaa	2160

Cys Pro Glu Gly Arg Pro Glu Phe Ser Glu Val Val Met Lys Leu Glu 705 710 715 720	
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agt ggg tct ctc tca cct tct tct tct tct gat tgc ctg gtg aac cgg Ser Gly Ser Leu Ser Pro Ser Ser Ser Ser Asp Cys Leu Val Asn Arg 740 745 750	2256
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Met Gly Asn Tyr Lys Ser Arg Pro Thr Gln Thr Cys Ser Asp Glu Trp
                  5
                                     10
aag aag aaa gtt agt gaa tot tac gct att atc ata gaa agg ctg gag
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Lys Lys Lys Val Ser Glu Ser Tyr Ala Ile Ile Ile Glu Arg Leu Glu
gat aac ctg cag atc aaa gaa aat gaa ttt caa gaa cta agg cac atc
                                                                   204
Asp Asn Leu Gln Ile Lys Glu Asn Glu Phe Gln Glu Leu Arq His Ile
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Phe Gly Ser Asp Glu Ala Phe Ser Glu Val Ser Leu Asn Tyr Arg Thr
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Glu Arg Gly Leu Ser Leu Leu His Leu Cys Cys Val Cys Gly Gly Asn
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aag tca cat atc cgt gcc ctt atg tta aaa ggg ctc cgt cca tcc aqa
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Lys Ser His Ile Arg Ala Leu Met Leu Lys Gly Leu Arg Pro Ser Arg
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ctg acg aga aat ggg ttt cca gct ctg cac ctg gcc gtt tac aag gac
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Leu Thr Arg Asn Gly Phe Pro Ala Leu His Leu Ala Val Tyr Lys Asp
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age ceg gaa ett ate act tea etg ttg cae age gga gea gat gtt cag
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Ser Pro Glu Leu Ile Thr Ser Leu Leu His Ser Gly Ala Asp Val Gln
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caa gtg gga tac ggt ggc ctc aca gcc ctc cac ata gct gca ata gct
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Gln Val Gly Tyr Gly Gly Leu Thr Ala Leu His Ile Ala Ala Ile Ala
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Gly His Pro Glu Ala Ala Glu Val Leu Leu Gln His Gly Ala Asn Val
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Asn Val Gln Asp Ala Val Phe Phe Thr Pro Leu His Ile Ala Ala Tyr
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					gct Ala 230											780
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					cag Gln											876
cct Pro	ttg Leu	cac His 275	ctg Leu	gca Ala	tgc Cys	tac Tyr	aat Asn 280	gga Gly	aat Asn	ttt Phe	gaa Glu	gtt Val 285	gcc Ala	aag Lys	gaa Glu	924
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ctg Leu	gtc Val	aaa Lys	ttt Phe	ctt Leu 325	ctt Leu	gat Asp	cag Gln	aat Asn	gct Ala 330	gtg Val	aac Asn	att Ile	aac Asn	cac His 335	cga Arg	1068
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tgt Cys 385	ttg Leu	atg Met	tgg Trp	gct Ala	tac Tyr 390	gag Glu	aaa Lys	gga Gly	cat His	gat Asp 395	gcc Ala	att Ile	gtt Val	aca Thr	ctc Leu 400	1260

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	-	_				_							_	atc Ile		1452
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tcc Ser	atc Ile 690	ccc Pro	aag Lys	ccc Pro	atc Ile	tca Ser 695	tcc Ser	ctg Leu	ctg Leu	ata Ile	cgg Arg 700	ggc Gly	tgg Trp	aat Asn	gca Ala	2172
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Phe Gly Ser Asp Glu Ala Phe Ser Glu Val Ser Leu Asn Tyr Arg Thr 50 55 60

Glu Arg Gly Leu Ser Leu Leu His Leu Cys Cys Val Cys Gly Gly Asn 65 70 75 80

Lys Ser His Ile Arg Ala Leu Met Leu Lys Gly Leu Arg Pro Ser Arg 85 90 95

Leu Thr Arg Asn Gly Phe Pro Ala Leu His Leu Ala Val Tyr Lys Asp 100 105 110

Ser Pro Glu Leu Ile Thr Ser Leu Leu His Ser Gly Ala Asp Val Gln 115 120 125

Gln Val Gly Tyr Gly Gly Leu Thr Ala Leu His Ile Ala Ala Ile Ala 130 135 140

Gly His Pro Glu Ala Ala Glu Val Leu Leu Gln His Gly Ala Asn Val 145 150 155 160

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Val Asn Val Ser Gly Glu Val Gly Asp Arg Pro Leu His Leu Ala Ser 195 200 205

Ala Lys Gly Phe Phe Asn Ile Val Lys Leu Leu Val Glu Glu Gly Ser 210 220

Lys Ala Asp Val Asn Ala Gln Asp Asn Glu Asp His Val Pro Leu His 225 230 235 240

Phe Cys Ser Arg Phe Gly His His Asn Ile Val Ser Tyr Leu Leu Gln 245 250 255

Ser Asp Leu Glu Val Gln Pro His Val Ile Asn Ile Tyr Gly Asp Thr 260 265 270

Pro Leu His Leu Ala Cys Tyr Asn Gly Asn Phe Glu Val Ala Lys Glu 275 280 285

Ile Val Gln Val Thr Gly Thr Glu Ser Leu Thr Lys Glu Asn Ile Phe 290 295 300

Ser Glu Thr Ala Phe His Ser Ala Cys Thr Tyr Gly Lys Asn Ile Asp 305 310 315 320

Leu Val Lys Phe Leu Leu Asp Gln Asn Ala Val Asn Ile Asn His Arg
325 330 335

Gly Arg Asp Gly His Thr Gly Leu His Ser Ala Cys Tyr His Gly His 340 345 350

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Cys Leu Met Trp Ala Tyr Glu Lys Gly His Asp Ala Ile Val Thr Leu 385 390 395 400

Leu Lys His Tyr Lys Arg Pro Gln Glu Glu Leu Pro Cys Asn Glu Tyr 405 410 415

Ser Gln Pro Gly Gly Asp Gly Ser Tyr Val Ser Val Pro Ser Pro Leu 420 425 430

Gly Lys Ile Lys Ser Met Thr Lys Glu Lys Ala Asp Val Leu Leu Leu 435

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Phe His Glu Ile Ile Gly Ser Gly Ser Phe Gly Lys Val Tyr Lys Gly

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Gln	Ser	Lys	Leu	Ile 565	Ile	Ala	Val	Asp	Val 570	Ala	Lys	Gly	Met	Glu 575	Tyr
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Glu	Ser 610	Arg	Phe	Leu	Gln	Ser 615	Leu	Asp	Glu	Asp	Asn 620	Met	Thr	Lys	Gln
Pro 625	Gly	Asn	Leu	Arg	Trp 630	Met	Ala	Pro	Glu	Val 635	Phe	Thr	Gln	Cys	Thr
Arg	Tyr	Thr	Ile	Lys 645	Ala	Asp	Val	Phe	Ser 650	Tyr	Ser	Leu	Cys	Leu 655	Trp
Glu	Leu	Leu	Thr 660	Gly	Glu	Ile	Pro	Phe 665	Ala	His	Leu	Lys	Pro 670	Ala	Ala
Ala	Ala	Ala 675	Asp	Met	Ala	Tyr	His 680	His	Ile	Arg	Pro	Pro 685	Ile	Gly	Tyr
Ser	Ile 690	Pro	Lys	Pro	Ile	Ser 695		Leu	Leu	Ile	Arg 700	_	Trp	Asn	Ala
Сув 705	Pro	Glu	Gly	Arg	Pro 710	Glu	Phe	Ser	Glu	Val 715	Val	Ser	Lys	Leu	Glu 720
Glu	Cys	Leu	Cys	Asn 725	Val	Glu	Leu	Met	Ser 730	Pro	Ala	Ser	Ser	Asn 735	Ser
Ser	Gly	Ser	Leu 740	Ser	Pro	Ser	Ser	Ser 745	Ser	Asp	Суз	Leu	Leu 750	Ser	Arg
Gly	Gly	Pro 755	Gly	Arg	Ser	His	Val 760	Ala	Ala	Leu	Arg	Ser 765	Arg	Phe	Glu

Leu Glu Tyr Ala Leu Asn Ala Arg Ser Tyr Ala Gly Trp Ser Gln Ser
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Val Gly Thr His Ser Asn Pro Gly Leu Ser Leu Glu Glu Met Asn Arg
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Gln Ser Lys Leu Ile Ile Ala Val Asp Val Ala Lys Gly Met Glu Tyr

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gac ag Asp Se	_	n									2505